



Input Set: N:\AMC\US08452843D.raw

Output Set: N:\CRF4\12052003\H452843D.raw

```
1 <110> APPLICANT: Sette, Alessandro
             Sidney, John
      3 <120> TITLE OF INVENTION: HLA Binding Peptides and Their Uses
      4 <130> FILE REFERENCE: 2060.0010002
C--> 5 <140> CURRENT APPLICATION NUMBER: US/08/452,843D
      6 <141> CURRENT FILING DATE: 1995-08-30
      7 <150> PRIOR APPLICATION NUMBER: US 08/344,824
      8 <151> PRIOR FILING DATE: 1994-11-23
      9 <150> PRIOR APPLICATION NUMBER: US 08/278,634
     10 <151> PRIOR FILING DATE: 1994-07-21
     11 <160> NUMBER OF SEQ ID NOS: 34
    12 <170> SOFTWARE: PatentIn ver 3.1
    14 <210> SEQ ID NO: 1
    15 <211> LENGTH: 9
    16 <212> TYPE: PRT
    17 <213> ORGANISM: Artificial Sequence
    18 <220> FEATURE:
    19 <223> OTHER INFORMATION: B35 consensus peptide
    20 <400> SEQUENCE: 1
              Phe Pro Phe Lys Tyr Ala Ala Ala Phe
                               5
    24 <210> SEQ ID NO: 2
    25 <211> LENGTH: 9
    26 <212> TYPE: PRT
    27 <213> ORGANISM: Artificial Sequence
    28 <220> FEATURE:
    29 <223> OTHER INFORMATION: Y1 analog of 1054.05
    30 <400> SEQUENCE: 2
    31
              Tyr Pro Lys Val Lys Gln Trp Pro Leu
    32
              1
    34 <210> SEQ ID NO: 3
    35 <211> LENGTH: 11
    36 <212> TYPE: PRT
    37 <213> ORGANISM: Artificial Sequence
    38 <220> FEATURE:
    39 <223> OTHER INFORMATION: MAGE-1
    40 <400> SEQUENCE: 3
    41
             Cys Ile Leu Glu Ser Cys Phe Arg Ala Val Ile
    44 <210> SEQ ID NO: 4
    45 <211> LENGTH: 9
    46 <212> TYPE: PRT
    47 <213> ORGANISM: Artificial Sequence
```

ENTERED

Input Set : N:\AMC\US08452843D.raw
Output Set: N:\CRF4\12052003\H452843D.raw

```
48 <220> FEATURE:
49 <223> OTHER INFORMATION: B53 self peptide
50 <400> SEQUENCE: 4
51
         Tyr Pro Ala Glu Ile Thr Leu Tyr Trp
52
54 <210> SEQ ID NO: 5
55 <211> LENGTH: 9
56 <212> TYPE: PRT
57 <213> ORGANISM: Artificial Sequence
58 <220> FEATURE:
59 <223> OTHER INFORMATION: Cw3 consensus
60 <400> SEQUENCE: 5
61
         Phe Ala Met Pro Asn Phe Gln Thr Leu
62
64 <210> SEQ ID NO: 6
65 <211> LENGTH: 9
66 <212> TYPE: PRT
67 <213> ORGANISM: Artificial Sequence
68 <220> FEATURE:
69 <223> OTHER INFORMATION: Cw3 consensus
70 <400> SEQUENCE: 6
71
         Phe Ala Met Pro Asn Phe Tyr Thr Leu
72
          1
74 <210> SEQ ID NO: 7
75 <211> LENGTH: 9
76 <212> TYPE: PRT
77 <213> ORGANISM: Artificial Sequence
78 <220> FEATURE:
79 <223> OTHER INFORMATION: Cw4 consensus
80 <400> SEQUENCE: 7
         Gln Pro Asp Asp Ala Val Tyr Lys Leu
81
84 <210> SEO ID NO: 8
85 <211> LENGTH: 9
86 <212> TYPE: PRT
87 <213> ORGANISM: Artificial Sequence
88 <220> FEATURE:
89 <223> OTHER INFORMATION: Cw6 consensus
90 <400> SEQUENCE: 8
91
         Ile Pro Tyr Pro Ile Val Arg Lys Leu
92
          1
94 <210> SEQ ID NO: 9
95 <211> LENGTH: 9
96 <212> TYPE: PRT
97 <213> ORGANISM: Artificial Sequence
98 <220> FEATURE:
99 <223> OTHER INFORMATION: Cw6 consensus
100 <400> SEQUENCE: 9
101
          Ile Pro Tyr Pro Ile Val Arg Ser Leu
```

Input Set: N:\AMC\US08452843D.raw
Output Set: N:\CRF4\12052003\H452843D.raw

```
102
104 <210> SEQ ID NO: 10
105 <211> LENGTH: 9
106 <212> TYPE: PRT
107 <213> ORGANISM: Artificial Sequence
108 <220> FEATURE:
109 <223> OTHER INFORMATION: Cw6 consensus
110 <400> SEQUENCE: 10
          Ile Pro Phe Pro Ile Val Arg Tyr Leu
111
112
                           5
          1
114 <210> SEQ ID NO: 11
115 <211> LENGTH: 9
116 <212> TYPE: PRT
117 <213> ORGANISM: Artificial Sequence
118 <220> FEATURE:
119 <223> OTHER INFORMATION: Histone H3.3
120 <400> SEQUENCE: 11
         Arg Tyr Arg Pro Gly Thr Val Ala Leu
122
          1
124 <210> SEQ ID NO: 12
125 <211> LENGTH: 9
126 <212> TYPE: PRT
127 <213> ORGANISM: Artificial Sequence
128 <220> FEATURE:
129 <223> OTHER INFORMATION: B7 Nat. Processed
130 <400> SEQUENCE: 12
131
          Met Pro Arg Gly Val Val Val Thr Leu
132
134 <210> SEQ ID NO: 13
135 <211> LENGTH: 10
136 <212> TYPE: PRT
137 <213> ORGANISM: Artificial Sequence
138 <220> FEATURE:
139 <223> OTHER INFORMATION: p53, 26-35
140 <400> SEQUENCE: 13
         Leu Pro Glu Asn Asn Val Leu Ser Pro Leu
141
                           5
144 <210> SEQ ID NO: 14
145 <211> LENGTH: 10
146 <212> TYPE: PRT
147 <213> ORGANISM: Artificial Sequence
148 <220> FEATURE:
149 <223> OTHER INFORMATION: p53, 84-93
150 <400> SEQUENCE: 14
151
         Ala Pro Ala Pro Ala Pro Ser Trp Pro Leu
152
                           5
154 <210> SEQ ID NO: 15
155 <211> LENGTH: 11
156 <212> TYPE: PRT
```

Input Set: N:\AMC\US08452843D.raw
Output Set: N:\CRF4\12052003\H452843D.raw

```
157 <213> ORGANISM: Artificial Sequence
158 <220> FEATURE:
159 <223> OTHER INFORMATION: p53, 127-137
160 <400> SEQUENCE: 15
         Ser Pro Ala Leu Asn Lys Met Phe Cys Gln Leu
                           5
164 <210> SEQ ID NO: 16
165 <211> LENGTH: 9
166 <212> TYPE: PRT
167 <213> ORGANISM: Artificial Sequence
168 <220> FEATURE:
169 <223> OTHER INFORMATION: p53, 154-162
170 <400> SEQUENCE: 16
         Gly Thr Arg Val Arg Ala Met Ala Ile
172
          1
174 <210> SEQ ID NO: 17
175 <211> LENGTH: 9
176 <212> TYPE: PRT
177 <213> ORGANISM: Artificial Sequence
178 <220> FEATURE:
179 <223> OTHER INFORMATION: p53, 249-257
180 <400> SEQUENCE: 17
         Arg Pro Ile Leu Thr Ile Ile Thr Leu
181
182
          1
184 <210> SEQ ID NO: 18
185 <211> LENGTH: 10
186 <212> TYPE: PRT
187 <213> ORGANISM: Artificial Sequence
188 <220> FEATURE:
189 <223> OTHER INFORMATION: p53, 299-308
190 <400> SEQUENCE: 18
191
      Leu Pro Pro Gly Ser Thr Lys Arg Ala Leu
192
         1 ·
                           5
194 <210> SEQ ID NO: 19
195 <211> LENGTH: 9
196 <212> TYPE: PRT
197 <213> ORGANISM: Artificial Sequence
198 <220> FEATURE:
199 <223> OTHER INFORMATION: p53, 315-323
200 <400> SEQUENCE: 19
201
         Ser Pro Gln Pro Lys Lys Pro Leu
202
          1
                           5
204 <210> SEQ ID NO: 20
205 <211> LENGTH: 10
206 <212> TYPE: PRT
207 <213> ORGANISM: Artificial Sequence
208 <220> FEATURE:
209 <223> OTHER INFORMATION: p53, 321-330
210 <400> SEQUENCE: 20
```

Input Set: N:\AMC\US08452843D.raw
Output Set: N:\CRF4\12052003\H452843D.raw

```
Lys Pro Leu Asp Gly Glu Tyr Phe Thr Leu
     211
     212
     214 <210> SEQ ID NO: 21
     215 <211> LENGTH: 9
     216 <212> TYPE: PRT
     217 <213> ORGANISM: Artificial Sequence
     218 <220> FEATURE:
     219 <223> OTHER INFORMATION: p53, 361-369
     220 <400> SEQUENCE: 21
               Gly Ser Arg Ala His Ser Ser His Leu
     221
     222
                                5
                1
     224 <210> SEQ ID NO: 22
     225 <400> SEQUENCE: 22
W--> 226
               000
     228 <210> SEQ ID NO: 23
     229 <400> SEQUENCE: 23
W--> 230
               000
     232 <210> SEQ ID NO: 24
     233 <400> SEQUENCE: 24
W--> 234
               000
     236 <210> SEQ ID NO: 25
     237 <400> SEQUENCE: 25
W--> 238
               000
     240 <210> SEQ ID NO: 26
     241 <211> LENGTH: 9
     242 <212> TYPE: PRT
     243 <213> ORGANISM: Artificial Sequence
     244 <220> FEATURE:
     245 <223> OTHER INFORMATION: supermotif
     246 <220> FEATURE:
     247 <221> NAME/KEY: VARIANT
     248 <222> LOCATION: (1)...(9)
     249 <223> OTHER INFORMATION: Xaa may be any of the amino acids found in Table 1, as
     250
               well as, non-naturally occurring amino acids or
     251
               amino acid mimetics
     252 <400> SEQUENCE: 26
W--> 253
               Xaa Pro Xaa Xaa Xaa Xaa Xaa Ala
     254
                                5
                1
     256 <210> SEO ID NO: 27
     257 <211> LENGTH: 9
     258 <212> TYPE: PRT
     259 <213> ORGANISM: Artificial Sequence
     260 <220> FEATURE:
     261 <223> OTHER INFORMATION: supermotif
     262 <220> FEATURE:
     263 <221> NAME/KEY: VARIANT
     264 <222> LOCATION: (1)...(9)
     265 <223> OTHER INFORMATION: Xaa may be any of the amino acids found in Table 1, as
               well as, non-naturally occurring amino acids or
     266
```

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/08/452,843D

DATE: 12/05/2003 TIME: 09:32:42

Input Set : N:\AMC\US08452843D.raw

Output Set: N:\CRF4\12052003\H452843D.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:26; Xaa Pos. 1,3,4,5,6,7,8 Seq#:27; Xaa Pos. 1,3,4,5,6,7,8 Seq#:28; Xaa Pos. 1,3,4,5,6,7,8 Seq#:29; Xaa Pos. 1,3,4,5,6,7,8 Seq#:30; Xaa Pos. 1,3,4,5,6,7,8 Seq#:31; Xaa Pos. 1,3,4,5,6,7,8 Seq#:32; Xaa Pos. 1,3,4,5,6,7,8,9 Seq#:33; Xaa Pos. 1,3,4,5,6,7,8,9,10 Seq#:34; Xaa Pos. 1,3,4,5,6,7,8,9,10,11